I hereby certify that this paper (along with any paper referred to as being attached or enclosed) is being deposited with the U.S. Postal Service on the date shown below with sufficient postage as First Class Mail, in an envelope addressed to: MS Amendment, Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450.

Dated: September 28, 2006 Signature:

(Maria Laccotripe Zacharakis, Ph.D., J.D.)

Docket No.: BGI-125CPCN (PATENT)

4 2006 E

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of:

Application No.: 10/627476

Confirmation No.: 2236

Filed: July 25, 2003

Art Unit: 1631

For:

CORYNEBACTERIUM GLUTAMICUM

VED.

Examiner: J. S. Brusca

GENES ENCODING PROTEINS INVOLVED IN MEMBRANE SYNTHESIS AND

MEMBRANE TRANSPORT

MS Amendment Commissioner for Patents P.O. Box 1450 Alexandria, VA 22313-1450

INFORMATION DISCLOSURE STATEMENT (IDS)

Dear Sir:

Pursuant to 37 CFR 1.56, 1.97 and 1.98, the attention of the Patent and Trademark Office is hereby directed to the references listed on the attached PTO/SB/08. It is respectfully requested that the information be expressly considered during the prosecution of this application, and that the references be made of record therein and appear among the "References Cited" on any patent to issue therefrom.

This Information Disclosure Statement is filed more than three months after the U.S. filing date, after the mailing date of the first Office Action on the merits, but before the mailing date of a Final Office Action or Notice of Allowance (37 CFR 1.97(c)).

For the Examiner's convenience in reviewing this continuation application, Applicants submit a consolidated PTO Form SB/08, listing all references cited during the prosecution of the parent application. References B29-B31 and C353-C355 were previously cited and submitted to the Office in U.S. Application Serial No. 09/602,787, filed June 23, 2000 (Atty. Docket No.

180.00 DA

Application No.: 10/627476 Docket No.: BGI-125CPCN

BGI-125CP), of which this application is a continuation and to which this application claims priority. In accordance with 37 CFR §1.98(d), these references are not enclosed herewith, but will be provided upon request. In addition, this Information Disclosure Statement presents references B1-B28 and C1-C352, and C356-357, not previously submitted to the Office, which are enclosed herewith in accordance with 37 CFR §1.97.

In accordance with 37 CFR 1.97(g), the filing of this Information Disclosure Statement shall not be construed to mean that a search has been made or that no other material information as defined in 37 CFR 1.56(a) exists. In accordance with 37 CFR 1.97(h), the filing of this Information Disclosure Statement shall not be construed to be an admission that any patent, publication or other information referred to therein is "prior art" for this invention unless specifically designated as such.

It is submitted that the Information Disclosure Statement is in compliance with 37 CFR 1.98 and the Examiner is respectfully requested to consider the listed references.

Please charge our Deposit Account No. 12-0080 in the amount of \$180.00 covering the fee set forth in 37 CFR 1.17(p). The Director is hereby authorized to charge any deficiency in the fees filed, asserted to be filed or which should have been filed herewith (or with any paper hereafter filed in this application by this firm) to our Deposit Account No. 12-0080, under Order No. BGI-125CPCN.

Dated: September 28, 2006

MLZ/MG/JGSmch

Respectfully submitted,

By Maria Laccotripe Zacharakis, Ph.D., J.D.

Registration No.: 56,266

LAHIVE & COCKFIELD, LLP

28 State Street

Boston, Massachusetts 02109

(617) 227-7400

(617) 742-4214 (Fax)

Attorney/Agent For Applicants

TRANSMITTAL

FORM (to be used for all correspondence after initial filing)

Total Number of Pages in This Submission

Serie to a composion of thiormasor	Torridge R displays a valid Civib control Hamber
Application Number	10/627476-Conf. #2236
Filing Date	July 25, 2003
First Named Inventor	Markus POMPEJUS
Art Unit	1631
Examiner Name	J. S. Brusca
Attorney Docket Number	BGI-125CPCN

	ENCLOSURES (Check all that apply)					
X Fee Transi	mittal Form	Drawing(s)		After Allowance Communication to TC		
Fee /	Attached	Licensing-related Papers		Appeal Communication to Board of Appeals and Interferences		
Amendmer	nt/Reply	Petition	•	Appeal Communication to TC (Appeal Notice, Brief, Reply Brief)		
After	Final	Petition to Convert to a Provisional Application		Proprietary Information		
Affida	avits/declaration(s)	Power of Attorney, Revocation Change of Correspondence		Status Letter		
Extension	of Time Request	Terminal Disclaimer		X Other Enclosure(s) (please Identify below):		
Express At	bandonment Request	Request for Refund		PTO form SB/08; Certificate of Mailing; Copies of Three Hundred		
x Information	n Disclosure Statement	CD, Number of CD(s)		and Eighty Two (382) References; Return Receipt Postcard		
Certified C	opy of Priority (s)	Landscape Table on	CD			
	lissing Parts/ Application	Remarks				
	y to Missing Parts under FR 1.52 or 1.53					
	1101.52 01 1.50					
	SIGNATI	URE OF APPLICANT, ATTOR	RNEY, OR	AGENT		
Firm Name	LAHIVE & COCKFIELD, KLP					
Signature		aous	M	1		
Printed name	Maria Laccotripe Za	charakis, Ph.D., J.D.		/		
Date	September 28, 2006		Reg. No.	56,266		

date shown below with sufficient postage as		g attriched or enclosed) is being deposited with the U.S. Postal Service on telope addressed to: MS Amendment, Commissioner for Patents, P.O. Box	he
1450, Alexandria, VA 22313-1450. Dated: September 28, 2006	Signature:	Maria Lactotripe Zagharakid, Ph.D., J.	ÐΛ

PTO/SB/17 (12-04v2) Approved for use through 7/31/2006. OMB 0651-0032 U.S. Patent and Trademark Office; U.S. DEPARTMENT OF COMMERCE Under the Paperwork Reduction Act of 1995, no person are required to respond to a collection of information unless it displays a valid OMB control number. Complete if Known Effective on 12/08/2004. 10/627476-Conf. #2236 Fees pursuant to the Consolidated Appropriations Act, 2005 (H.R. 4818). **Application Number** FEE TRANSMITTAL July 25, 2003 Filing Date Markus POMPEJUS First Named Inventor For FY 2005 **Examiner Name** J. S. Brusca Applicant claims small entity status. See 37 CFR 1.27 1631 Art Unit **BGI-125CPCN** TOTAL AMOUNT OF PAYMENT (\$) 180.00 Attorney Docket No. METHOD OF PAYMENT (check all that apply) Credit Card None Check Money Order Other (please identify): Deposit Account Number: 12-0080 Deposit Account Name: Lahive & Cockfield, LLP For the above-identified deposit account, the Director is hereby authorized to: (check all that apply) Charge fee(s) indicated below Charge fee(s) indicated below, except for the filing fee Charge any additional fee(s) or underpayment of Credit any overpayments fee(s) under 37 CFR 1.16 and 1.17 **FEE CALCULATION** 1. BASIC FILING, SEARCH, AND EXAMINATION FEES **FILING FEES** SEARCH FEES **EXAMINATION FEES** Small Entity Small Entity **Small Entity Application Type** Fee (\$) Fee (\$) Fee (\$) Fee (\$) Fees Paid (\$) Fee (\$) Fee (\$) Utility 300 150 500 250 200 100 200 100 100 50 130 65 Design 200 100 300 150 80 Plant 160 300 150 500 250 600 Reissue 300 200 100 0 0 **Provisional** 0 0 2. EXCESS CLAIM FEES **Small Entity** Fee (\$) Fee (\$) Fee Description Each claim over 20 (including Reissues) 50 25 Each independent claim over 3 (including Reissues) 200 100 Multiple dependent claims 360 180 **Multiple Dependent Claims** Fee Paid (\$) **Total Claims Extra Claims** Fee (\$) Fee Paid (\$) Fee (\$) Fee Paid (\$) Indep. Claims **Extra Claims** Fee (\$) 3. APPLICATION SIZE FEE If the specification and drawings exceed 100 sheets of paper (excluding electronically filed sequence or computer listings under 37 CFR 1.52(e)), the application size fee due is \$250 (\$125 for small entity) for each additional 50 sheets or fraction thereof. See 35 U.S.C. 41(a)(1)(G) and 37 CFR 1.16(s). Number of each additional 50 or fraction thereof Fee Paid (\$) **Total Sheets** Extra Sheets Fee (\$) - 100 = _____/50 (round up to a whole number) x 4. OTHER FEE(S) Fees Paid (\$) Non-English Specification, \$130 fee (no small entity discount) Other (e.g., late filing surcharge): 1806 Submission of an Information Disclosure Statement 180.00 SUBMITTED BY Registration No. (Altorney/Agent) 56,266 Telephone (617) 227-7400 Signature

date shown below with sufficient	llong with any paper re postage as First Class	erred to as being attached or enclosed) is being depos Mail, in an envelope addressed to: MS Amendment,	sited with the U.S. Postal Service on the Commissioner for Patents, P.O. Box 1450,
Alexandria, VA 22313-1450.			
		IX AAAAAA	24
D 0 -1 -1 00 0000	Simulation /	account Maria Laccount	Achadia Dh.D. ID.
Dated: September 28, 2006	Signature:	(Mana Caccoth	pe //acharakis, Ph.D., J.D.)

Date

September 28, 2006

Dated: September 28, 2006

Name (Print/Type)

Maria Laccotripe Zacharakis, Ph.D., J.D.

PTO/SB/08a/b (07-05)

Approved for use through 07/31/2006. OMB 0651-0031

U.S. Patent and Trademark Office; U.S. DEPARTMENT OF COMMERCE Under the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it contains a valid OMB control nu

Substitute for form 1449A/B/PTO

INFORMATION DISCLOSURE STATEMENT BY APPLICANT

(Use as many sheets as necessary)

19 1 Sheet of

Complete if Known					
Application Number	10/627476-Conf. #2236				
Filing Date	July 25, 2003				
First Named Inventor Markus POMPEJUS					
Art Unit	1631				
Examiner Name	John S. Brusca				
Attorney Docket Number	BGI-125CPCN				

U.S. PATENT DOCUMENTS							
Examiner Initials*	Cite No. ¹	Document Number Number-Kind Code ² (if known)	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear		

		FOREI	GN PATENT	DOCUMENTS		
Examiner Initials*	Cite No. ¹	Foreign Patent Document Country Code ³ -Number ⁴ -Kind Code ⁵ (if known)	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear	Te
	B1	EP-0204326-A2	12-10-1986	Kyowa Hakko Kogyo Co., Ltd.		
	B2	JP-62232392	10-12-1987	Kyowa Hakko Kogyo Co., Ltd.		Abstr.
	B3	JP-62244382	10-24-1987	Ajinomoto Co., Inc.		Abstr.
	B4	EP-0358940-A1	03-21-1990	Degussa Aktiengesellschaft		
	B5	JP-04278088	10-02-1992	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B6	JP-04330284	11-18-1992	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B7	JP-05030977	02-09-1993	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B8	JP-05056782	03-09-1993	Kyowa Hakko Kogyo Co. Ltd.		Abstr.
	B9	JP-05076352	03-30-1993	Ajinomoto Co., Inc.		Abstr.
	B10	JP-05184366	07-27-1993	Mitsubishi Petrochem Co. Ltd.		Abstr.
	B11	JP-05184371	07-27-1993	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B12	JP-05284970	11-02-1993	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B13	JP-05284972	11-02-1993	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B14	JP-05344881	12-27-1993	Ajinomoto Co., Inc.		Abstr.
	B15	JP-05344893	12-27-1993	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B16	JP-06062866	03-08-1994	Ajinomoto Co., Inc.		Abstr.
	B17	JP-06169780 /	06-21-1994	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B18	JP-06261766 /	09-20-1994	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B19	JP-06277067	10-04-1994	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B20	JP-06277073	10-04-1994	Mitsubishi Petrochem Co., Ltd.		Abstr.
	B21	JP-07031476	02-03-1995	Mitsubishi Chem	-	Abstr.
	B22	JP-07031478	02-03-1995	Mitsubishi Chem		Abstr.
	B23	JP-09028391	02-04-1997	Mitsubishi Chem		Abstr.
	B24	JP-09070291	03-18-1997	Ajinomoto Co., Inc.		Abstr.
	B25	JP-07075578	03-20-1995	Mitsubishi Chem		Abstr.
	B26	JP-07075579	03-20-1995	Mitsubishi Chem		Abstr.
	B27	WO-9519442-A1	07-20-1995	Forschungszentrum Jülich GMGH Möckel		Abstr.
	B28	JP-09224661	09-02-1997	Mitsubishi Chem		Abstr.
	B29*	EP-0252558-A2		SCLAVO S.p.A.		
		EP-0752472-B1		Ajinomoto Co. Inc.		
		EP-0786519-A2		Human Genome Sciences, Inc.		

*EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant. Applicant's unique citation designation number (optional). See Kinds Codes of USPTO Patent Documents at <u>www.uspto.gov</u> or MPEP 901.04. ³ Enter Office that issued the document, by the two-letter code (WIPO Standard ST.3). ⁴ For Japanese patent documents, the indication of the year of the reign of the Emperor must precede the serial number of the patent document. ⁵ Kind of document by the appropriate symbols as indicated on the document under WIPO Standard ST.16 if possible. Applicant is to place a check mark here if English language Translation is attached.

Examiner	Date	
Signature	Considered	

stitute for form 1449A/B/	PTO		Complete if Known		
			Application Number	10/627476-Conf. #2236	
FORMATIC	N DISC	CLOSURE	Filing Date	July 25, 2003	
TATEMENT	BY AP	PLICANT	First Named Inventor	Markus POMPEJUS	
			Art Unit	1631	
(Use as many sheets as necessary)			Examiner Name	John S. Brusca	
2	of	19	Attorney Docket Number	BGI-125CPCN	
	FORMATIC TATEMENT	(Use as many sheets as nee	FORMATION DISCLOSURE TATEMENT BY APPLICANT (Use as many sheets as necessary)	FIRST Named Inventor (Use as many sheets as necessary) Application Number Filing Date First Named Inventor Art Unit Examiner Name	

		NON PATENT LITERATURE DOCUMENTS	
Examiner Initials	Cite No. ¹	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T ²
	C1	Ankri, Serge, et al., "Mutations in the Corynebacterium glutamicum Proline Biosynthetic Pathway: a Natural Bypass of the proA Step," Journal of Bacteriology, Vol. 178(15):4412-4419 (1996)	
	C2	Billman-Jacobe, H., "Nucleotide sequence of a recA gene from Comybacterium glutamicum," The Journal of Sequencing and Mapping," Vol. 4:403-404 (1994)	
	C3	Bonamy, Celine, et al., "Identification of IS1206, a Corynebacterium glutamicum IS3-related insertion sequence and phylogenetic analysis," Molecular Microbiology, Vol. 14(3):571-581 (1994)	
	C4	Bonnassie, S., et al., "Nucleotide sequence of the dapA gene from Corynebacterium glutamicum," Nucleic Acids Research, Vol. 18(21):6421 (1990)	
	C5	Börmann, E.R., et al., "Molecular analysis of the Corynebacterium glutamicum gdh gene encoding glutamate dehydrogenase," Molecular Microbiology, Vol. 6(3):317-326 (1992)	
	C6	Chen, Chian-Chi, et al., "The cloning and nucleotide sequence of a Corynebacterium glutamicum 3-deoxy-D-arabinoheptulosonate-7-phosphate synthase gene," FEMS Micrkobiology Letters, Vol. 107:223-230 (1993)	
	C7	Cianciotto, Nicholas, et al., "DNA sequence homology between attB-related sites of Corynebacterium diphtheriae, Corynebacterium ulcerans, Corynebacterium glutamicum, and the attP site of γ-Corynephage," FEMS Microbiology Letters, Vol. 66:299-302 (1990)	
	C8	Correia, Antonio, et al., "Cloning and characterization of an IS-like element present in the genome of <i>Brevibacterium lactofermentum</i> ATCC 13869," <i>Gene,</i> Vol. 170:91-94 (1996)	
	C9	Dusch, Nicole, et al., "Expression of the Corynebacterium glutamicum panD Gene Encoding L-Aspartate-α-Decarboxylase Leads to Pantothenate Overproduction in Escherichia coli," Applied and Environmental Microbiology, Vol. 65(4):1530-1539 (1999)	
	C10	Eikmanns, Bernhard J., et al., "Nucleotide sequence, expression and transcriptional analysis of the <i>Corynebacterium glutamicum gltA</i> gene encoding citrate synthase," <i>Microbiology</i> , Vol. 140:1817-1828 (1994)	
	C11	Eikmanns, Bemhard J., "Identification, Sequence Analysis, and Expression of a Corynebacterium glutamicum Gene Cluster Encoding the Three Glycolytic Enzymes Glyceraldehyde-3-Phosphate Dehydrogenase, 3-Phosphoglycerate Kinase, and Triosephosphate Isomerase," Journal of Bacteriology, Vol. 174(19):6076-6086 (1992)	
	C12	Eikmanns, Bemhard J., et al., "Cloning, Sequence Analysis, Expression, and Inactivation of the Corynebacterium glutamicum icd Gene Encoding Isocitrate Dehydrogenase and Biochemical Characterization of the Enzyme," Journal of Bacteriology, Vol. 177(3):774-782 (1995)	
	C13	Eikmanns, Bernhard J., et al., "The phosphoenolpyruvate carboxylase gene of Corynebacterium glutamicum: Molecular cloning, nucleotide sequence, and expression," Mol. Gen. Genet., Vol. 218:330-339 (1989)	
	C14	Fitzpatrick, R., et al., "Construction and characterization of recA mutant strains of Corynebacterium glutamicum and Brevibacterium lactofermentum," Appl. Microbiol. Biotechnol., Vol. 42:575-580 (1994)	
	C15	Follettie, Max T., et al., "Molecular Cloning and Nucleotide Sequence of the Corynebacterium glutamicum pheA Gene," Journal of Bacteriology, Vol. 167(2):695-702 (1986)	
	C16	Fouet, Agnes, et al., "Bacillus subtilis sucrose-specific enzyme II of the phosphotransferase system: Expression in Escherichia coli and homology to enzymes II from enteric bacteria,"	

Examiner		"	Date	
Signature			Considered	

Sut	Substitute for form 1449A/B/PTO			Complete if Known		
				Application Number	10/627476-Conf. #2236	
11	NFORMATION	I DIS	SCLOSURE	Filing Date	July 25, 2003	
S	TATEMENT I	BY A	PPLICANT	First Named Inventor	Markus POMPEJUS	
}				Art Unit	1631	
	(Use as many sheets as necessary)			Examiner Name	John S. Brusca	
Sheet	3	of	19	Attorney Docket Number	BGI-125CPCN	

	Proc. Natl. Acad. Sci., Vol. 84:8773-8777 (1987)
C17	Han, KS., et al., "The molecular structure of the Corynebacterium glutamicum threonine synthase gene," Molecular Microbiology, Vol. 4(10):1693-1702 (1990)
C18	Heery, D.M., et al., "Nucleotide sequence of the Corynebacterium glutamicum trpE gene," Nucleic Acids Research, Vol. 18(23):7138 (1990)
C19	Heery, D.M., et al., "Cloning of the trp Gene Cluster from a Tryptophan-Hyperproducing Strain of Corynebacterium glutamicum: Identification of a Mutation in the trp Leader Sequence," Applied and Environmental Microbiology, Vol. 59(3):791-799 (1993)
C20	Heery, David M., et al., "A Sequence from a Tryptophan-Hyperproducing Strain of Corynebacterium glutamicum Encoding Resistance to 5-Methyltryptophan," Biochemcial and Biophysical Research Communications, Vol. 201(3):1255-1262 (1994)
C21	Honrubia, M.P., et al., "Identification, characterization, and chromosomal organization of the ftsZ gene from Brevibacterium lactofermentum," Mol. Gen. Genet., Vol. 259:97-104 (1998)
C22	Ishino, Shuichi, et al., "Nucleotide sequence of the meso-diaminopimelate D-dehydrogenase gene from Corynebacterium glutamicum," <i>Nucleic Acids Research,</i> Vol. 15(9):3917 (1987)
C23	Jäger, Wolfgang, et al., "A Corynebacterium glutamicum Gene Conferring Multidrug Resistance in the Heterologous Host Escherichia coli," Journal of Bacteriology, Vol. 179(7):2449-2451 (1997)
C24	Jäger, Wolfgang, et al., "A Corynebacterium glutamicum gene encoding a two-domain protein similar to biotin carboxylases and biotin-carboxyl-carrier proteins," Arch. Microbiol., Vol. 166:76-82 (1996)
C25	Jakoby, Marc, et al., "Isolation of the Corynebacterium glutamicum glnA gene encoding glutamine synthetase I," FEMS Microbiology Letters, Vol. 154:81-88 (1997)
C26	Jakoby, Marc, et al., "Nitrogen regulation in Corynebacterium glutamicum: isolation of genes involved and biochemical characterization of corresponding proteins," FEMS Microbiology Letters, Vol. 173:303-310 (1999)
C27	Jetten, Mike S., et al., "Structural and Functional Analysis of Pyruvate Kinase from Corynebacterium glutamicum," Applied and Environmental Microbiology, Vol. 60(7):2501-2507 (1994)
C28	Joliff, G., et al., "Cloning and nucleotide sequence of the csp1 gene encoding PS1, one of the two major secreted proteins of Corynebacterium glutamicum: the deduced N-terminal region of PS1 is similar to the Mycobacterium antigen 85 complex," Molecular Microbiology, Vol. 6(16):2349-2362 (1992)
C29	Kalinowski, J., et al., "Genetic and biochemical analysis of the aspartokinase from Corynebacterium glutamicum," Molecular Microbiology, Vol. 5(5):1197-1204 (1991)
C30	Kalinowski, Jörn, et al., "Aspartokinase gene $lysC_{\chi}$ and $lysC_{\beta}$ overlap and are adjacent to the aspartate β-semialdehyde dehydrogenase gene asd in Corynebacterium glutamicum," Mol. Gen. Genet., Vol. 224:317-324 (1990)
C31	Keilhauer, Carmen, et al., "Isoleucine Synthesis in Corynebacterium glutamicum: Molecular Analysis of the ilvB-ilvN-ilvC Operon," Journal of Bacteriology, Vol. 175(17):5595-5603 (1993)
C32	Kimura, Eiichiro, et al., "Molecular Cloning of a Novel Gene, dtsR, Which Rescues the Detergent Sensitivity of a Mutant Derived from Brevibacterium lactofermentum," Biosci. Biotech. Biochem., Vol. 60(10):1565-1570 (1996)
C33	Kobayashi, Miki, et al., "Cloning, Sequencing, and Characterization of the ftsZ Gene from Coryneform Bacteria," Biochemical and Biophysical Research Communications, Vol. 236:383-388 (1997)
C34	Kronemeyer, Wolfgang, et al., "Structure of the <i>gluABCD</i> Cluster Encoding the Glutamate Uptake System of Corynebacterium glutamicum," Journal of Bacteriology, Vol. 177(5):1152-1158 (1995)

			· · · · · · · · · · · · · · · · · · ·
Examiner		Date	
Signature	1 ·	Considered	

Sub	ostitute for form 1449A/B/PT	o			Complete if Known
				Application Number	10/627476-Conf. #2236
	NFORMATION	1 DI	SCLOSURE	Filing Date	July 25, 2003
S	STATEMENT BY APPLICANT			First Named Inventor	Markus POMPEJUS
				Art Unit	1631
	(Use as many sheets as necessary)		Examiner Name	John S. Brusca	
Sheet	4	of	19	Attorney Docket Number	BGI-125CPCN

C35	Lee, Heung-Shick, et al., "Molecular Characterization of AceB, a Gene Encoding Malate Synthase in Corynebacterium glutamicum," Journal of Microbiology and Biotechnology, Vol. 4(4):256-263 (1994)	
C36	Le Marrec, Claire, et al., "Genetic Characterization of Site-Specific Integration Functions of ΦAAU2 Infecting 'Arthrobacter aureus' C70," Journal of Bacteriology, Vol. 178(7):1996-2004 (1996)	
C37	Lepiniec, Loïc, et al., "Sorghum phosphoenolpyruvate carboxylase gene family: structure, function and molecular evolution," Plant Molecular Biology, Vol. 21:487-502 (1993)	
C38	Lichtinger, Thomas, et al., "Biochemical and Biophysical Characterization of the Cell Wall Porin of Corynebacterium glutamicum: The Channel Is Formed by a Low Molecular Mass Polypeptide," Biochemistry, Vol. 37:15024-15032 (1998)	
C39	Ludwig, W., et al., "Phylogenetic relationships of <i>Bacteria</i> based on comparative sequence analysis of elongation factor Tu and ATP-synthase β-subunit genes," <i>Antonie van Leeuwenhoek</i> , Vol. 64:285-305 (1993)	
C40	Malubres, Marcos, et al., "Analysis said Expression of the thrC Gene of Brevibacterium lactofermentum and Characterization of the Encoded Threonine Synthase," Applied and Environmental Microbiology, Vol. 60(7):2209-2219 (1994)	
C41	Marcel, T., et al., "Nucleotide sequence and organization of the upstream region of the Corynebacterium glutamicum lysA gene," Molecular Microbiology, Vol. 4(11):1819-1830 (1990)	
C42	Mateos, Luis M., et al., "Nucleotide sequence of the homoserine kinase (thr B) gene of Brevibacterium lactofermentum," Nucleic Acids Research, Vol. 15(9):3922 (1987)	
C43	Mateos, Luis M., et al., "Nucleotide sequence of the homoserine dehydrogenase (thr A) gene of Brevibacterium lactofermentum," Nucleic Acids Research, Vol. 15(24):10598 (1987)	
C44	Matsui, Kazuhiko, et al., "Complete nucleotide and deduced amino acid sequences of the Brevibacterium lactofermentum tryptophan operon," Nucleic Acids Research, Vol. 14(24):10113-10114 (1986)	
C45	Möckel, Bettina, et al., "Functional and Structural Analyses of Threonine Dehydratase from Corynebacterium glutamicum," Journal of Bacteriology, Vol. 174(24):8065-8072 (1992)	
C46	Molenaar, Douwe, et al., "Biochemical and genetic characterization of the membrane-associated malate dehydrogenase (acceptor) from Corynebacterium glutamicum," Eur. J. Biochem., Vol. 254:395-403 (1998)	
C47	Moreau, Sylvia, et al., "Site-specific integration of corynephage Φ16: construction of an integration vector," <i>Microbiology</i> , Vol. 145:539-548 (1999)	
C48	Moreau, Sylvie, et al., "Analysis of the Integration Functions of Φ304L: An Integrase Module among Corynephages," <i>Virology</i> , Vol. 255:150-159 (1999)	
C49	O'Gara, James P., et al., "Mutations in the <i>trpD</i> Gene of <i>Corynebacterium glutamicum</i> Confer 5-Methyltryptophan Resistance by Encoding a Feedback-Resistant Anthranilate Phosphoribosyltransferase," <i>Applied and Environmental Microbiology</i> , Vol. 61(12):4477-4479 (1995)	
C50	Oguiza, José A., et al., "A Gene Encoding Arginyl-tRNA Synthetase Is Located in the Upstream Region of the IysA Gene in <i>Brevibacterium lactofermentum</i> : Regulation of <i>argS-lysA</i> Cluster Expression by Arginine," <i>Journal of Bacteriology</i> , Vol. 175(22):7356-7362 (1993)	
C51	Oguiza, José A., et al., "Molecular Cloning, DNA Sequence Analysis, and Characterization of the Corynebacterium diphtheriae dtxR Homolog from Brevibacterium lactofermentum," Journal of Bacteriology, Vol. 177(2):465-467 (1995)	
C52	Oguiza, José A., et al., "Multiple Sigma Factor Genes in <i>Brevibacterium lactofermentum</i> : Characterization of sigA and sigB," Journal of Bacteriology, Vol. 178(2):550-553 (1996)	

Examiner	Date
Signature	Considered

Sut	ostitute for form 1449A/B	/PTO	*		Complete if Known
				Application Number	10/627476-Conf. #2236
11	NFORMATIC	ON DIS	CLOSURE	Filing Date	July 25, 2003
S	STATEMENT BY APPLICANT			First Named Inventor	Markus POMPEJUS
				Art Unit	1631
	(Use as many	sheets as n	ecessary)	Examiner Name	John S. Brusca
Sheet	5	of	19	Attorney Docket Number	BGI-125CPCN

C53	Oguiza, José A., et al., "The galE gene encoding the UDP-galactose 4-epimerase of	—
	Brevibacterium lactofermentum is coupled transcriptionally to the dmdR gene," Gene, Vol. 177:103-107 (1996)	
C54	O'Regan, Michael, et al., "Cloning and nucleotide sequence of the phosphoenolpyruvate carboxylase-coding gene of <i>Corynebacterium glutamicum</i> ATCC13032," <i>Gene</i> , Vol. 77:237-251 (1989)	
C55	Park, Soo-Dong, et al., "Isolation and Analysis of metA, a Methionine Biosynthetic Gene Encoding Homoserine Acetyltransferase in Corynebacterium glutamicum," Mol. Cells, Vol. 8(3):286-294 (1998)	
C56	Park, Yong-Ha, et al., "Phylogenetic Analysis of the Coryneform Bacteria by 5S rRNA Sequences," <i>Journal of Bacteriology,</i> Vol. 169(5):1801-1806 (1987)	
C57	Pascual, Cristina, et al., "Phylogenetic Analysis of the Genus Corynebacterium Based on 16S rRNA Gene Sequences," International Journal of Systematic Bacteriology, Vol. 45(4):724-728 (1995)	****
C58	Pátek, M., et al., "Analysis of the leuB gene from Corynebacterium glutamicum," Appl Microbiol Biotechnol, Vol. 50:42-47 (1998)	
C59	Pátek, Miroslav, et al., "Leucine Synthesis in Corynebacterium glutamicum: Enzyme Activities, Structure of leuA, and Effect of leuA Inactivation on Lysine Synthesis," Applied and Environmental Microbiology, Vol. 60(1):133-140 (1994)	
C60	Pátek, Miroslav, et al., "Promoters from Corynebacterium glutamicum: cloning, molecular analysis and search for a consensus motif," Microbiology, Vol. 142:1297-1309 (1996)	
C61	Pátèk, M., et al., "Identification and transcriptional analysis of the dapB-ORF2-dapA-ORF4 operon of Corynebacterium glutamicum, encoding two enzymes involved in L-lysine synthesis," Biotechnology Letters, Vol. 19(11):1113-1117 (1997)	
C62	Peoples, O.P., et al., "Nucleotide sequence and fine structural analysis of the Corynebacterium glutamicum hom-thrB operon," Molecular Microbiology, Vol. 2(1):63-72 (1988)	
C63	Peter, Heidi, et al., "Corynebacterium glutamicum Is Equpped with Four Secondary Carriers for Compatible Solutes: Identification, Sequencing, and Characterization of the Proline/Ectoine Uptake System, ProP, and the Ectoine/Proline/Glycine Betaine Carrier, EctP," Journal of Bacteriology, Vol. 180(22):6005-6012 (1998)	
C64	Peter, Heidi, et al., "Isolation of the <i>putP</i> gene of <i>Corynebacterium glutamicum</i> and characterization of a low-affinity uptake system for compatible solutes," <i>Arch Microbiol</i> , Vol. 168:143-151 (1997)	
C65	Peter, Heidi, et al., "Isolation, Characterization, and Expression of the Corynebacterium glutamicum betP Gene, Encoding the Transport System for the Compatible Solute Glycine Betaine," Journal of Bacteriology, Vol. 178(17):5229-5234 (1996)	
C66	Peters-Wendisch, Petra G., et al., "Pyruvate carboxylase from Corynebacterium glutamicum: characterization, expression and inactivation of the pyc gene," Microbiology, Vol. 144:915-927 (1998)	
C67	Peyret, J.L., et al., "Characterization of the cspB gene encoding PS2, an ordered surface-layer protein in Corynebacterium glutamicum," Molecular Biology, Vol. 9(1):97-109 (1993)	
C68	Pisabarro, Agustín, et al., "A Cluster of Three Genes (dapA, orf2, and dapB) of Brevibacterium lactofermentum Encodes Dihydrodipicolinate Synthase, Dihydrodipicolinate Reductase, and a Third Polypeptide of Unknown Function," Journal of Bacteriology, Vol. 175(9):2743-2749 (1993)	
C69	Rainey, Frederick A., et al., "Phylogenetic analysis of the genera <i>Rhodococcus</i> and <i>Nocardia</i> and evidence for the evolutionary origin of the genus <i>Nocardia</i> from within the radiation of <i>Rhodococcus</i> species," <i>Microbiology</i> , Vol. 141:523-528 (1995)	

Examiner	 Date	
Signature	Considered	1

Sub	estitute for form 1449A/B/PT	О			Complete if Known
				Application Number	10/627476-Conf. #2236
10	NFORMATION	1 DI	SCLOSURE	Filing Date	July 25, 2003
S	TATEMENT B	3Y A	APPLICANT	First Named Inventor	Markus POMPEJUS
				Art Unit	1631
	(Use as many sheets as necessary)			Examiner Name	John S. Brusca
Sheet	6	of	19	Attorney Docket Number	BGI-125CPCN

C70	Ramos, Adoración, et al., "Cloning, sequencing and expression of the gene encoding elongation factor P in the amino-acid producer <i>Brevibacterium lactofermentum</i> (Corynebacterium glutamicum ATCC 13869)," Gene, Vol. 198:217-222 (1997)	
C71	Reinscheid, Dieter, J., et al., "Cloning, sequence analysis, expression and inactivation of the Corynebacterium glutamicum pta-ack operon encoding phosphotransacetylase and acetate kinase," Microbiology, Vol. 145:503-513 (1999)	
C72	Reinscheid, Dieter J., et al., "Characterization of the Isocitrate Lyase Gene from Corynebacterium glutamicum and Biochemical Analysis of the Enzyme," Journal of Bacteriology, Vol. 176(12):3474-3483 (1994)	
C73	Reinscheid, Dieter J., et al., "Malate synthase from Corynebacterium glutamicum: sequence analysis of the gene and biochemical characterization of the enzyme," <i>Microbiology</i> , Vol. 140:3099-3108 (1994)	
C74	Roller, Carsten, et al., "Gram-positive bacteria with a high DNA G+C content are characterized by a common insertion within their 23S rRNA genes," <i>Journal of General Microbiology</i> , Vol. 138:1167-1175 (1992)	
C75	Rossol, Ingrid, et al., "The Corynebacterium glutamicum aecD Gene Encodes a C-S Lyase with α,β-Elimination Activity That Degrades Aminoethylcysteine," <i>Journal of Bacteriology</i> , Vol. 174(9):2968-2977 (1992)	
C76	Ruimy, Raymond, et al., "Phylogeny of the Genus Corynebacterium Deduced from Analyses of Small-Subunit Ribosomal DNA Sequences," <i>International Journal of Systematic Bacteriology</i> , Vol. 45(4):740-746 (1995)	
C77	Sahm, Hermann, et al., "D-Pantothenate Synthesis in Corynebacterium glutamicum and Use of panBC and Genes Encoding L-Valine Synthesis for D-Pantothenate Overproduction," Applied and Environmental Microbiology, Vol. 65(5):1973-1979 (1999)	
C78	Sakanyan, Vehary, et al., "Genes and enzymes of the acetyl cycle of arginine biosynthesis in Corynebacterium glutamicum: enzyme evolution in the early steps of the arginine pathway," Microbiology, Vol. 142:99-108 (1996)	
C79	Sano, Konosuke, et al., "Structure and function of the <i>trp</i> operon control regions of Brevibacterium lactofermentum, a glutamic-acid-producing bacterium," Gene, Vol. 53:191-200 (1987)	
C80	Schäfer, Andreas, et al., "Cloning and Characterization of a DNA Region Encoding a Stress-Sensitive Restriction System from Corynebacterium glutamicum ATCC 13032 and Analysis of Its Role in Intergeneric Conjugation with Escherichia coli," Journal of Bacteriology, Vol. 176(23):7309-7319 (1994)	
C81	Seep-Feldhous, A.J., et al., "Molecular analysis of the Corynebacterium glutamicum lysl gene involved in lysine uptake," Molecular Microbiology, Vol. 5(12):2995-3005 (1991)	
C82	Serebrijski, I, et al., "Multicopy Suppression by asd Gene and Osmotic Stress-Dependent Complementation by Heterologous proA in proA Mutants," Journal of Bacteriology, Vol. 177(24):7255-7260 (1995)	
C83	Serebriiskii, Ilya G., et al., "Two new members of the BioB superfamily: cloning, sequencing and expression of bioB genes of Methylobacillus flagellatum and Corynebacterium glutamicum," Gene, Vol. 175:15-22 (1996)	
C84	Siewe, Ruth M. et al, "Functional and Genetic Characterization of the (Methyl)ammonium Uptake Carrier of Corynebacterium glutamicum," The Journal of Biological Chemistry, Vol. 271(10):5398-5403 (1996)	
C85	Usuda, Yoshihiro, et al., "Molecular cloning of the Corynebacterium glutamicum ('Brevibacterium lactofermentum' AJ12036) odhA gene encoding a novel type of 2-oxoglutarate dehydrogenase," Microbiology, Vol. 142:3347-3354 (1996)	
C86	Vertès, Alain A., et al., "Isolation and characterization of IS31831, a transposable element from Corynebacterium glutamicum," Molecular Microbiology, Vol. 11(4):739-746 (1994)	

Examiner	Date
Signature	Considered

BGI-125CPCN

PTO/SB/08a/b (07-05)

Approved for use through 07/31/2006. OMB 0651-0031

U.S. Patent and Trademark Office; U.S. DEPARTMENT OF COMMERCE

Under the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it contains a valid OMB control number. Complete if Known Substitute for form 1449A/B/PTO 10/627476-Conf. #2236 Application Number INFORMATION DISCLOSURE July 25, 2003 Filing Date STATEMENT BY APPLICANT Markus POMPEJUS First Named Inventor 1631 Art Unit (Use as many sheets as necessary) John S. Brusca Examiner Name

Attorney Docket Number

7

of

19

Sheet

	C87	von der Osten, C.H., et al., "Molecular cloning, nucleotide sequence and fine-structural analysis of the <i>Corynebacterium glutamicum fda</i> gene: structural comparison of <i>C. glutamicum</i> fructose-1,6-biphosphate aldolase to class I and class II aldolases," <i>Molecular Microbiology</i> , Vol. 33(11):1625-1637 (1989)
	C88	Vrljic, Marina, et al., "A new type of transporter with a new type of cellular function: L-lysine export from Corynebacterium glutamicum," Molecular Microbiology, Vol. 22(5):815-826 (1996)
	C89	Wachi, M., et al., "A <i>mur</i> C gene from coryneform bacteria," <i>Appl Microbiol Biotechnol</i> , Vol. 51:223-228 (1999)
	C90	Wehmeier, Lutz, et al., "The role of the Corynebacterium glutamicum rel gene in (p)ppGpp metabolism," Microbiology, Vol. 144:1853-1862 (1998)
	C91	Wehrmann, Axel, et al., "Different Modes of Diaminopimelate Synthesis and Their Role in Cell Wall Integrity: a Study with Corynebacterium glutamicum," Journal of Bacteriology, Vol. 180(12):3159-3165 (1998)
	C92	Wehrmann, Axel, et al., "Analysis of different DNA fragments of Corynebacterium glutamicum complementing dapE of Escherichia coli," Microbiology, Vol. 140:3349-3356 (1994)
	C93	Wehrmann, Axel, et al., "Functional Analysis of Sequences Adjacent to dapE of Corynebacterium glutamicum Reveals the Presence of aroP, Which Encodes the Aromatic Amino Acid Transporter," Journal of Bacteriology, Vol. 177(20):5991-5993 (1995)
	C94	Yeh, Patrice, et al., "Nucleotide sequence of the <i>lysA</i> gene of <i>Corynebacterium glutamicum</i> and possible mechanisms for modulation of its expression," <i>Mol Gen Genet</i> , Vol. 212:112-119 (1988)
	C95	GenBank Accession No. A09073 for DNA fragment coding for phosphoenolpyruvat corboxylase, recombinant DNA carrying said fragment, strains carrying the recombinant DNA and method for producing L-amino acids using said strains, Bachmann, B. et al, 04/14/2005
	C96	GenBank Accession No. A45579 for Production of L-isoleucine by means of recombinant micro-organisms with deregulated threonine dehydratase, Moeckel, B. et al, 03/07/1997
	C97	GenBank Accession No. A45581, Moeckel, B. et al, "Production of L-Isoleucine by Means of Recombinant Micro-Organisms with Deregulated Threonine Dehydratase." 03/07/97
	C98	GenBank Accession No. A45583, Moeckel, B. et al, "Production of L-Isoleucine by Means of Recombinant Micro-Organisms with Deregulated Threonine Dehydratase." 03/07/97
	C99	GenBank Accession No. A45585, Moeckel, B. et al, "Production of L-Isoleucine by Means of Recombinant Micro-Organisms with Deregulated Threonine Dehydratase." 03/07/97
	C100	GenBank Accession No. A45587 for Production of L-isoleucine by means of recombinant micro-organisms with deregulated threonine dehydratase, Moeckel, B. et al, 03/07/97
	C101	GenBank Accession No. AA011641 for Generation and analysis of 280,000 human expressed sequence tags, Hillier, L. et al, 05/09/97
	C102	GenBank Accession No. AA655226 for The WashU-HHMI Mouse EST Project, Marra, M. et al, 11/04/97
	C103	GenBank Accession No. AA704727 for WashU-NCI human EST Project, Hillier, L. et al, 12/24/97
	C104	GenBank Accession No. AB003132 for Cloning, sequencing, and characterization of the ftsZ gene from coryneform bacteria, Kobayashi, M. et al, 08/04/97
	C105	GenBank Accession No. AB015023 for A murC gene from coryneform bacteria, Wachi, M. et al, 12/15/01
	C106	GenBank Accession No. AB015853 for Expression in Escherichia coli of a new multidrug efflux pump, MexXY, from Pseudomonas aeruginosa, Mine, T. et al, 08/02/00
	C107	GenBank Accession No. AB018530 for Molecular cloning of a novel gene, dtsR, which rescues the detergent sensitivity of a mutant derived from Brevibacterium lactofermentum, Kimura, E. et al, 10/16/98
	C108	GenBank Accession No. AB018531 for The role of DtsR2 in the glutamate-production in coryneform bacteria, Kimura, E. et al, 10/16/98
aminer Inature		Date Considered

BGI-125CPCN

PTO/SB/08a/b (07-05)

Approved for use through 07/31/2006. OMB 0651-0031

U.S. Patent and Trademark Office; U.S. DEPARTMENT OF COMMERCE

Under the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it contains a valid OMB control number. Complete if Known Substitute for form 1449A/B/PTO 10/627476-Conf. #2236 Application Number **INFORMATION DISCLOSURE** July 25, 2003 Filing Date STATEMENT BY APPLICANT Markus POMPEJUS First Named Inventor Art Unit 1631 (Use as many sheets as necessary) John S. Brusca **Examiner Name**

Attorney Docket Number

19

8

of

Sheet

T	0400	ConDonis According No. AD0000004 for the United States of the condition of						
1 .	C109	GenBank Accession No. AB020624 for Isolation of the murl gene from Brevibacterium lactofermentum ATCC 13869 encoding D-glutamate racemase, Malathi, K.C. et al, 07/24/99						
+	C110							
	CIIO	glutamicum transketolase gene, Ikeda, M. et al, 02/20/9						
	C111	GenBank Accession No. AB024708 for Corynebacterium glutamicum gltBD gene, Kanno, S. et al, 03/13/99						
	C112	GenBank Accession No. AB025424 for Brevibacterium lactofermentum ATCC 13869 acn gene for Aconitase, Nakamura, J. et al, 04/03/99						
	C113	GenBank Accession No. AB027714 for Cryptic plasmid pCG1 of Corynebacterium glutamicum, Yonetani, Y. et al, 06/01/99						
	C114	GenBank Accession No. AB027715 for Plasmid pCG11 of Corynebacterium glutamicum, Yonetani, Y. et al, 06/01/99						
	C115	GenBank Accession No. AC004295 for Sequencing of Drosophila chromosome 2R, region 55C1-55C4, Celniker, S.E. et al, 07/29/98						
	C116	GenBank Accession No. AC005019 for Toward a complete human genome sequence, Sulston, J.E. et al, 10/15/03						
	C117	GenBank Accession No. AC006044 for Toward a complete human genome sequence, Sulston, J.E. et al, 10/08/03						
	C118	GenBank Accession No. AC006474 for The DNA sequence of human chromosome 7, Hillier, L.W. et al, 01/27/04						
	C119	GenBank Accession No. AC007084 for Sequencing of Drosophila chromosome 2R, region 43F-44A, Celniker, S.E. et al, 03/21/01						
	C120	GenBank Accession No. AC007739 for The sequence of Homo sapiens BAC clone RP11- 91L3, Hou, S. et al, 04/16/05						
	C121	GenBank Accession No. AC008403, DOE Joint Genome Institute and Stanford Human Genome Center, for Homo sapiens chromosome 19 clone CTC-273B12, complete sequence, 03/22/03						
	C122	GenBank Accession No. AC009298 for The sequence of Homo sapiens BAC clone RP11-1716, Nguyen, C. et al, 04/15/05						
	C123	GenBank Accession No. AC010187 for Homo sapiens chromosome 12 clone RP11-389O9, Working Draft Sequence, 40 unordered pieces, Muzny, D.M. et al, 01/08/03						
	C124	GenBank Accession No. AC011647 for Homo sapiens chromosome 11, clone RP11-15D18, Birren, B. et al, 06/18/02						
	C125	GenBank Accession No. AD000016 for Mycobacterium tuberculosis, Du, L., 12/03/96						
-	C126	GenBank Accession No. AD000002 for Mycobacterium tuberculosis, Du, L., 12/10/96						
	C127	GenBank Accession No. AE000654 for The complete genome sequence of the gastric pathogen Helicobacter pylori, Tomb, JF. et al, 04/06/99						
	C128	GenBank Accession No. AF001552 for Genome duplications and other features in 12 Mb of DNA sequence from human chromosome 16p and 16q, Loftus, B.J. et al, 01/01/00						
	C129	GenBank Accession No. AF005242, Chun, J.Y. et al, "Molecular cloning and analysis of the argC gene from Corynebacterium glutamicum," <i>Biochem. Mol. Biol. Int.</i> , Vol. 46(3):437-447 (1998), 07/23/01						
	C130	GenBank Accession No. AF005635 for Corynebacterium glutamicum, Reid, S.J. et al, 06/14/99						
	C131	GenBank Accession No. AF030405 for Molecular cloning of the histidine biosynthetic genes from Corynebacterium glutamicum, Jung, S.I. et al, 11/13/97						
	C132	GenBank Accession No. AF030520, Ko, S.Y. et al, "Molecular cloning of the argG gene from Corynebacterium glutamicum." 11/19/97						
	C133	GenBank Accession No. AF031518, Chun, J.Y. et al, "Cloning of the argF gene encoding the ornithine carbamoyltransferase from Corynebacterium glutamicum," <i>Mol. Cells</i> , Vol. 9(3):333-337 (1999), 06/13/01						
Examiner	1	Date						
Signature	1	Considered						

Sub	stitute for form 1449A/B/PT	го		Complete if Known		
				Application Number	10/627476-Conf. #2236	
16	IFORMATION	A DI	SCLOSURE	Filing Date	July 25, 2003	
S	TATEMENT I	BY F	APPLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many sh	ieets as	necessary)	Examiner Name	John S. Brusca	
Sheet	9	of	19	Attorney Docket Number	BGI-125CPCN	

	C134	GenBank Accession No. AF036932 for Molecular cloning of the aroD gene from
		Corynebacterium glutamicum, Park, KY. et al, 12/13/97
	C135	GenBank Accession No. AF038548 for Sequence of the Corynebacterium glutamicum pyruvate carboxylase gene, Koffas, M.A. et al, 08/25/00
	C136	GenBank Accession No. AF038651 for The role of the Corynebacterium glutamicum rel gene
		in (p)ppGpp metabolism, Wehmeier, L. et al, 03/15/01
(C137	GenBank Accession No. AF041436, Ko. SY. et al, "The argR gene of Corynebacterium
	0.100	glutamicum." 01/05/99
I '	C138	GenBank Accession No. AF045998 for Molecular cloning of the histidine biosynthetic genes
		from Corynebacterium glutamicum, Jund, S.I. et al, 02/19/98
•	C139	GenBank Accession No. AF048764, Park. M.Y. et al, "Molecular cloning of the argH gene
		encoding argininosuccinate lyase from Corynebacterium glutamicum." 07/01/98
	C140	GenBank Accession No. AF049897, Park, M.Y. et al, "Molecular cloning of the Arginine Biosynthetic Gene from Corynebacterium glutamicum." 07/01/98
	C141	GenBank Accession No. AF050109 for The function of the inhA gene in mycolic acid synthesis
		of Corynebacterium glutamicum, Sayyada-Hafeez, A. et al, 11/19/02
	C142	GenBank Accession No. AF050166, Kwon, J.H. et al, "Cloning of the histidine biosynthetic
		genes from Corynebacterium glutamicum: organization and analysis of the hisG and hisE
		genes," Can. J. Microbiol., Vol. 46(9):848-855 (2000), 04/26/02
[C143	GenBank Accession No. AF051846 for Corynebacterium glutamicum, Jung, S.I. et al, 03/12/98
	C144	GenBank Accession No. AF052652 for Isolation and Analysis of metA, a Metionine
		Biosynthetic Gene Encoding Homoserine Acetyltransferase in Corynebacterium glutamicum,
		Park, SD. et al, 03/19/98
	C145	GenBank Accession No. AF053071 for Cloning and analysis of the aroB gene encoding
		dehyddroquinate synthase from Corynebacterium glutamicum, Han, M.A. et al, 04/26/02
(C146	GenBank Accession No. AF060558 for Corynebacterium glutamicum, Juns, S.I. et al, 04/29/98
	C147	GenBank Accession No. AF077409, Latreille, P. et al, "The A. thaliana Genome Sequencing
		Project," August 3, 1998
1	C148	GenBank Accession No. AF086704 for Corynebacterium glutamicum, Kwon, J.H. et al,
		02/08/99
	C149	GenBank Accession No. AF114233 for Cloning and molecular analysis of the
		Corynebacterium glutamicum ASO19 and aroA gene, O'Donohue, M. et al, 02/07/99
	C150	GenBank Accession No. AF116184, Dusch, N. et al, "Expression of the Corynebacterium
		glutamicum panD gene encoding L-aspartate-alpha-decarboxylase leads to pantothenate
		overproduction in Escherichia coli," Appl. Environ. Microbiol., Vol. 65(4):1530-1539 (1999),
		05/02/99
	C151	GenBank Accession No. AF124518 for The cloning and phylogenetic analysis of the 3-
		dehydroquinase gene from Corynebacterium glutamicum, Joy, J. et al, 05/18/99
Į•	C152	GenBank Accession No. AF124600 for Genetic aspects of the prechorismate pathway in
		Corynebacterium glutamicum, Burke, K.G. et al, 05/04/99
- 10	C153	GenBank Accession No. AF145898 for Comparison of inhA gene of Corynebacterium
		glutamicum with its mutants and other related species, Hafeez, S. A. et al, 11/19/02
	C154	GenBank Accession No. AF145897 for Comparison of inhA gene for Corynebacterium
		glutamicum with its mutants and other related species, Hafeez, S.A. et al, 11/19/02
10	C155	GenBank Accession No. Al190741, NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap, for
		National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index,
		10/28/98
	C156	GenBank Accession No. AJ001436 for Corynebacterium glutamicum is equipped with four
		secondary carriers for compatible solutes: identification, sequencing, and characterization of
		the proline/ectoine uptake system, ProP, and the ectoine/proline/glycine betaine carrier, EctP,
		Peter, H. et al, 11/20/98
aminer		Date
gnature		Considered
2	<u> </u>	Journalite

Sub	stitute for form 1449A/B/P	то		Complete if Known		
				Application Number	10/627476-Conf. #2236	
10	IFORMATION	V DISC	CLOSURE	Filing Date	July 25, 2003	
S	TATEMENT	BY AP	PLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many sh	eets as ned	essary)	Examiner Name	John S. Brusca	
Sheet	10	of	19	Attorney Docket Number	BGI-125CPCN	

[C157	GenBank Accession No. AJ004934, Wehrmann, A. et al, "Different modes of diaminopimelate synthesis and their role in cell wall integrity: a study with Corynebacterium glutamicum," J.
		Bacteriol., Vol. 180(12):3159-3165 (1998), 06/17/98
[1	C158	GenBank Accession No. AJ007732, Jakoby, M.J. et al, "Ammonium uptake in
		Corynebacterium glutamicum is regulated on the level of expression and enzyme activity." 04/15/05
	C159	GenBank Accession No. AJ010319 for Nitrogen regulation in Corynebacterium glutamicum:
		isolation of genes involved and biochemical characterization of corresponding proteins, Jakoby, M. et al, 04/15/05
10	C160	GenBank Accession No. AJ132968 for Construction and application of new Corynebacterium
- 1		glutamicum vectors, Jakoby, M.J. et al, 05/04/99
1	C161	GenBank Accession No. AJ224946 for Biochemical and genetic characterization of the
		membrane-associated malate dehydrogenase (acceptor) from Corynebacterium glutamicum,
1		Molenaar, D. et al, 08/11/98
	C162	GenBank Accession No. AJ238250 for Functions of the membrane-associated and
		cytoplasmic malate dehydrogenases in the citric acid cycle of Corynebacterium glutamicum,
1		Molenaar, D. et al, 12/03/00
	C163	GenBank Accession No. AJ238703 for Biochemical and biophysical characterization of the cell
1		wall porin of Corynebacterium glutamicum: the channel is formed by a low molecular mass
		polypeptide, Lichtinger, T. et al, 05/07/99
	C164	GenBank Accession No. AL021841, Cole, S.T. et al, "Deciphering the biology of
i		Mycobacterium tuberculosis from the complete genome sequence," Nature, Vol.
		393(6685):537-544 (1998), 09/02/02
[4	C165	GenBank Accession No. AL022075 for Deciphering the biology of Mycobacterium tuberculosis
		from the complete genome sequence, Cole, S.T. et al, 09/02/02
Į (C166	GenBank Accession No. AL022268 for A set of ordered cosmids and a detailed genetic and
8		physical map for the 8 Mb Streptomyces coelicolor A3 (2) chromosome, Redenbach, M. et al, 05/12/02
į.	C167	GenBank Accession No. AL023591 for Use of an ordered cosmid library to deduce the
		genomic organization of Mycobacterium leprae, Eiglmeier, K. et al, 04/16/05
	C168	GenBank Accession No. AL035159 for Use of an ordered cosmid library to deduce the
	···	genomic organization of Mycobacterium leprae, Eiglmeier, K. et al, 04/16/05
	C169	GenBank Accession No. AL081678 for Arabidopsis thaliana, Salanoubat, M. et al, 06/28/99
(C170	GenBank Accession No. AL096814 for Homo sapiens, Sehra, H., 05/18/05
	<u>C171</u>	GenBank Accession No. AL101527 for Drosophila melanogaster, Genoscope, 07/26/99
[1	C172	GenBank Accession No. AP000004 for Complete sequence and gene organization of the
		genome of a hyper-thermophilic archaebacterium, Pyrococcus horikoshii OT3, Kawarabayasi, Y. et al, 05/27/04
	C173	GenBank Accession No. AP000140 for Homo sapiens 911,949bp genomic DNA of 21q21.2
	0170	(Region: LL56-APP Clone Range: B2291C14-R44F3), Hattori, M. et al, 01/08/00
	C174	GenBank Accession No. AP000228 for Homo sapiens 75,698bp genomic DNA of 21q21.2,
	• • • •	Hattori, M. et al, 11/20/99
	C175	GenBank Accession No. AQ128685 for Sequence-tagged connectors: A sequence approach
		to mapping and scanning the human genome, Mahairas, G.G. et al, 09/23/98
	C176	GenBank Accession No. AQ364540 for A BAC End Sequencing Framework to Sequence the
		Rice Genome, Wing, R.A. et al, 12/16/99
1	C177	GenBank Accession No. AQ463737 for Sequence-tagged connectors: A sequence approach
		to mapping and scanning the human genome, Mahairas, G.G. et al, 04/23/99
- 1	C178	GenBank Accession No. B10133 for BAC End Sequences at ATGC, Feng, J. et al, 05/14/97
	C179	GenBank Accession No. C97772 for Rice cDNA from callus, Sasaki, T. et al, 04/04/02
	C173	
	C180	GenBank Accession No. D17429 for Isolation and characterization of IS31831, a transposable

Sub	estitute for form 1449A/B/PT	0		Complete if Known		
				Application Number	10/627476-Conf. #2236	
11	IFORMATION	DIS	CLOSURE	Filing Date	July 25, 2003	
S	TATEMENT E	BY A	PPLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many she	eets as i	necessary)	Examiner Name	John S. Brusca	
Sheet	11	of	19	Attorney Docket Number	BGI-125CPCN	

	element from Corynebacterium glutamicum, Vertes, A.A. et al, 02/04/99						
C181	GenBank Accession No. D84102 for Molecular cloning of the Corynebacterium glutamicum						
0.01	('Brevibacterium lactofermentum' AJ12036) odhA gene encoding a novel type of 2-						
i	oxoglutarate dehydrogenase, Usuda, Y. et al, 02/06/99						
C192							
C182	GenBank Accession No. D84432 for Systematic sequencing of the 283 kb 210 degrees-232						
	degrees region of the Bacillus subtilis genome containing the skin element and many						
	sporulation genes, Mizuno, M. et al, 02/06/99						
C183	GenBank Accession No. D87915 for Molecular cloning and characterization of the obg gene of						
	Streptomyces griseus in relation to the onset of morphological differentiation, Okamoto, S. et						
	al, 02/07/99						
C184	GenBank Accession No. E01358 for Production of L-Thereonine and L-Isoleucine, Katsumata,						
	R. et al, 09/29/97						
C185	GenBank Accession No. E01359 for Production of L-Thereonine and L-Isoleucine, Katsumata,						
0,00	R. et al, 09/29/97						
C186							
C100	GenBank Accession No. E01375 for Tryptophan operon, peptide and protein coded thereby,						
	utilization of Tryptophan operon gene expression and producation of Tryptophan, Matsui, K. et						
	al, 09/29/97						
C187	GenBank Accession No. E01376 for Tryptophan Operon, Peptide and Protein Coded Thereby,						
	Utilization of Tryptophan Operon Gene Expression and Production of Tryptophan, Matsui, K.						
	et al, 09/29/97						
C188	GenBank Accession No. E01377 for Tryptophan Operon, Peptide and Protein Coded Thereby,						
	Utilization of Tryptophan Operon Gene Expression and Production of Tryptophan, Matsui, K.						
	et al, 09/29/97						
C189	GenBank Accession No. E03937 for DNA fragment containing gene capable of coding biotin						
0.00	synthetase and its utilization, Hatakeyama, K. et al, 09/29/97						
C190	GenBank Accession No. E04040 for Gene Coding Diaminopelargonic Acid Aminotransferase						
0130	and Desthiobiotin Synthetase and its Utilization, Kohama, K. et al, 09/29/97						
C191							
Cigi	GenBank Accession No. E04041 for Gene coding Diaminopelargonic Acid Aminotransferase						
0400	and Desthiobiotin Synthetase and its Utilization, Kohama, K. et al, 09/29/97						
C192	GenBank Accession No. E04307, Kurusu, Y. et al, "Gene DNA coding aspartase and						
	utilization there." 09/29/97						
C193	GenBank Accession No. E04376 for Gene Manifestation Controlling DNA, K atsumata, R. et						
	al, 09/29/97						
C194	GenBank Accession No. E04377 for Gene Manifestation Controlling DNA, Katsumata, R. et al,						
	09/29/97						
C195	GenBank Accession No. E04484 for Production of L-Phenylalanine by Fermentation, Sotouchi,						
	N. et al, 09/29/97						
C196	GenBank Accession No. E05108 for Gene DNA Coding Aspartokinase and its Use, Fugono,						
	N. et al, 09/29/97						
C197	GenBank Accession No. E05112 for Gene DNA Coding Dihydrodipicolinic Acid Synthetase						
0.137	and its Use, Hatakeyama, K. et al, 09/29/97						
C109							
C198	GenBank Accession No. E05776 for Gene DNA Coding Diaminopimelic Acid Dehydrogenase						
0.155	and its Use, Kobayashi, M. et al, 09/29/97						
C199	GenBank Accession No. E05779 for Gene DNA Coding Threonine Synthase and its Use,						
	Kohama, K. et al, 09/29/97						
C200	GenBank Accession No. E06110 for Production of L-Phenylalanine by Fermentation Method,						
	Kikuchi, T. et al, 09/29/97						
C201	GenBank Accession No. E06111 for Production of L-Phenylalanine by Fermentation Method,						
0201	Kikuchi, T. et al, 09/29/97						
C202	GenBank Accession No. E06146 for Gene Capable of Coding Acetohydroxy Acid Synthase						

Examiner	Date	
Signature	Considered	

Sub	stitute for form 1449A/B/P	го		Complete if Known		
				Application Number	10/627476-Conf. #2236	
	IFORMATION	V DI	SCLOSURE	Filing Date	July 25, 2003	
S	TATEMENT	BY A	APPLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many sh	ieets as	s necessary)	Examiner Name	John S. Brusca	
Sheet	12	of	19	Attorney Docket Number	BGI-125CPCN	

	and its, Inui, M. et al, 09/29/97
C203	GenBank Accession No. E06825 for Mutant Aspartokinase Gene, Sugimoto, M. et al, 09/29/97
C204	GenBank Accession No. E06826 for Mutant Aspartokinase Gene, Sugimoto, M. et al, 09/29/97
C205	GenBank Accession No. E06827 for Mutant Aspartokinase Gene, Sugimoto, M. et al, 09/29/97
C206	GenBank Accession No. E07701 for Gene DNA Participating in Integration of Membraneous
	Protein to, Honno, N. et al, 09/29/97
C207	GenBank Accession No. E08177 for Genetic DNA Capable of Coding Aspartokinase Released
	from Feedback Inhibition and its Utilization, Sato, Y. et al, 09/29/97
C208	GenBank Accession No. E08178 for Genetic DNA Capable of Coding Aspartokinase Released
	from Feedback Inhibition and its Utilization, Sato, Y. et al, 09/29/97
 C209	GenBank Accession No. E08179 for Genetic DNA Capable of Coding Aspartokinase Released
	from Feedback Inhibition and its Utilization, Sato, Y. et al, 09/29/97
C210	GenBank Accession No. E08180 for Genetic DNA Capable of Coding Aspartokinase Released
	from Feedback Inhibition and its Utilization, Sato, Y. et al, 09/29/97
C211	GenBank Accession No. E08181 for Genetic DNA Capable of Coding Aspartokinase Released
	from Feedback Inhibition and its Utilization, Sato, Y. et al, 09/29/97
C212	GenBank Accession No. E08182 for Genetic DNA Capable of Coding Aspartokinase Released
	from Feedback Inhibition and its Utilization, Sato, Y. et al, 09/29/97
C213	GenBank Accession No. E08232 for Gene DNA Coding Acetohydroxy Acid Isomeroreductase,
	Inui, M. et al, 09/29/97
C214	GenBank Accession No. E08234 for Gene DNA Coding for Translocation Machinery of
	Protein, Asai, Y. et al, 09/29/97
C215	GenBank Accession No. E08643 for DNA Fragment Having Promoter Function in Coryneform
	Bacterium, Hatakeyama, K. et al, 09/29/97
C216	GenBank Accession No. E08646 for DNA Fragment Having Promoter Function in Coryneform
	Bacterium, Hatakeyama, K. et al, 09/29/97
C217	GenBank Accession No. E08649, Kohama, K. et al, "DNA fragment having promoter function
	in coryneform bacterium." 09/29/97
C218	GenBank Accession No. E08900 for DNA Fragment Containig Gene Coding Dihyrodipicolinic
	Acid Reductase and Utilization Thereof, Madori, M. et al, 09/29/97
C219	GenBank Accession No. E08901 for DNA Fragment Containing Gene Coding Dihyrodipicolinic
	Acid Decarboxylase and Utilization Thereof, Madori, M. et al, 09/29/97
C220	GenBank Accession No. E12594 for Production of L-Tryptophan, Hatakeyama, K. et al,
	09/29/97
C221	GenBank Accession No. E12758 for Amplification of Gene Using Artificial Transposon, Moriya,
	M. et al, 09/29/97
C222	GenBank Accession No. E12759 for Amplification of Gene Using Artificial Transposon, Moriya,
	M. et al, 09/29/97
C223	GenBank Accession No. E12760 for Amplification of Gene Using Artificial Transposon, Moriya,
	M. et al, 09/29/97
C224	GenBank Accession No. E12764 for Amplification of Gene Using Artificial Transposon, Moriya,
	M. et al, 09/29/97
C225	GenBank Accession No. E12767 for Amplification of Gene Using Artificial Transposon, Moriya,
	M. et al, 09/29/97
C226	GenBank Accession No. E12770 for Amplification of Gene Using Artificial Transposon, Moriya,
	M. et al, 09/29/97
 C227	GenBank Accession No. E12773 for Amplification of Gene Using Artificial Transposon, Moriya,
	M. et al, 09/29/97
 C228	GenBank Accession No. E13655 for Glucose-6-Phosphate Dehydrogenase and DNA Capable
	of Coding the Same, Hatakeyama, K. et al, 04/27/98

·		
Examiner	Date	
Signature	Considered	

Subs	stitute for form 1449A/B/	PTO		Complete if Known		
				Application Number	10/627476-Conf. #2236	
IN	IFORMATIC	on disc	CLOSURE	Filing Date	July 25, 2003	
S	TATEMENT	BYAP	PLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many	sheets as ned	cessary)	Examiner Name	John S. Brusca	
Sheet	13	of	19	Attorney Docket Number	BGI-125CPCN	

C220	ConPort Associan No. LO4509, Market B. et al. "Expetiend and structural and services of	
C229	,	
	threonine dehydratase from Corynebacterium glutamicum," <i>J. Bacteriol.</i> , Vol. 174(24):8065-	
	8072 (1992), 04/26/93	
C230	3	
	Corynebacterium glutamicum 3-deoxy-D-arabinoheptulosonate-7-phosphate synthase gene,	
	Chen, C.C. et al, 04/26/93	
C231	GenBank Accession No. L09232 for Isoleucine synthesis in Corynebacterium glutamicum:	
	molecular analysis of the ilvB-ilvN-ilvC operon, Keilhauer, C. et al, 02/23/95	
C232		•
	phosphotransferase system: expression in Escherichia coli and homology to enzymes II from	
ì	enteric bacteria, Fouet, A. et al, 11/24/94	
C233		
0200	malate synthase in Corynebacterium glutamicum, Lee, HS. et al, 06/08/95	
C234		
0204		
	kinase from Corynebacterium glutamicum," Appl. Environ. Microbiol., Vol. 60(7):2501-2507	
0005	(1994), 12/07/94	
C235		
	isocitrate lyase in corynebacterium glutamicum, Lee, HS et al, 02/10/95	
C236		
	and characterization of the Corynebacterium diphtheria dtxR homolog from Brevibacterium	
	lactofermentum," J. Bacteriol., Vol. 177(2):465-467 (1995) 03/06/96	
C237	GenBank Accession No. L78820, Eiglmeier, K. et al, "Use of an ordered cosmid library to	
	deduce the genomic organization of Mycobacterium laprae," Mol. Microbiol., Vol. 7(2):197-206	
	(1993) 12/17/01	
C238		
0200	sequence of the Corynebacterium glutamicum pheA gene," <i>J. Bacteriol.</i> , Vol. 167(2):695-702	
l i	(1986), 04/26/93	
C239	GenBank Accession No. M16175 for Phylogenetic analysis of the coryneform bacteria by 5S	
0200	rRNA sequences, Park, Y.H. et al, 04/27/93	
C240		
C240	GenBank Accession No. M16663 for Structure and function of the trp operon control regions of	
0044	Brevibacterium lactofermentum, a glutamic-acid producing bacterium, Sano, K. et al, 06/11/01	
C241	GenBank Accession No. M16664 for Structure and function of the trp operon control regions of	
	Brevibacterium lactofermentum, a glutamic-acid-producing bacterium, Sano, K. et al, 06/11/01	
C242	The state of the s	
	phosphoenolpyruvate carboxylase-coding gene of Corynebacterium glutamicum ATCC13032,	
	O'Regan, M. et al, 12/15/95	
C243	GenBank Accession No. M85106 for Gram-positive bacteria with a high DNA G+C content are	
	characterized by a common insertion within their 23S rRNA genes, Roller, C. et al, 04/26/93	
C244	GenBank Accession No. M85107 for Gram-positive bacteria with a high DNA G+C content are	
	characterized by a common insertion within their 23S rRNA genes, Roller, C. et al, 04/26/93	
C245	GenBank Accession No. M85108 for Gram-positive bacteria with a high DNA G+C content are	
	characterized by a common insertion within their 23S rRNA genes, Roller, C. et al, 04/26/93	
C246		
C240	GenBank Accession No. M89931, Rossol, I. et al, "The Corynebacterium glutamicum aecD	
	gene encodes a C-S lyase with alpha, beta-elimination activity that degrades	
	aminoethylcysteine," <i>J. Bacteriol.</i> , Vol. 174(9):2968-2977 (1992), 02/08/02	
C247	GenBank Accession No. S59299 for Cloning of the trp gene cluster from a tryptophan-	
	hyperproducing strain of Corynebacterium glutamicum: identification of a mutation in the trp	
	leader sequence, Herry, D.M. et al, 04/13/01	
C248	GenBank Accession No. S76966 for Insertion sequence typing of Mycobacterium tuberculosis:	
	characterization of a widespread subtype with a single copy of IS6110, Fomukong, N.G. et al,	
	05/11/05	
Evaminer	Date	

Examiner	Date		
Signature	Considered		

Substitute for form 1449A/B/PTO			Complete if Known
	Applio	cation Number	10/627476-Conf. #2236
Information disclos	SURE Filing	Date	July 25, 2003
STATEMENT BY APPLIC	CANT First I	Named Inventor	Markus POMPEJUS
	Art U	nit	1631
(Use as many sheets as necessary)	Exam	iner Name	John S. Brusca
Sheet 14 of	19 Attorr	ney Docket Number	BGI-125CPCN

C249	GenBank Accession No. U00016 for Mycobacterium leprae, Smith, D.R. et al, 03/01/04
C250	GenBank Accession No. U00018, Smith, D.R., Mycobacterium laprae, Smith, D.R. et al, 03/01/04
C251	GenBank Accession No. U11545 for Complete nucleotide sequence of the Corynebacterium glutamicum ATCC 21850 trpD gene, O'Gara, J.P. et al, 07/08/94
C252	GenBank Accession No. U13922 for Cloning and characterization of a DNA region encoding a stress-sensitive restriction system from Corynebacterium glutamicum ATCC 13032 and analysis of its role in intergeneric conjugation with Escherichia coli, Schafer, A. et al, 02/03/98
C253	GenBank Accession No. U14965 for Molecular cloning and characterization of the recA gene from Corynebacterium glutamicum ASO19, Kerins, S.M. et al, 02/05/99
C254	GenBank Accession No. U31224, Ankri, S. et al, "Mutations in the Corynebacterium glutamicum proline biosynthetic pathway: a natural bypass of the proA step," <i>J. Bacteriol.</i> , Vol. 178(15):4412-4419 (1996), 08/02/96
C255	GenBank Accession No. U31225, Ankri, S. et al, "Mutations in the Corynebacterium glutamicum proline biosynthetic pathway: a natural bypass of th proA step," <i>J. Bacteriol.</i> , Vol. 178(15):4412-4419 (1996), 08/02/96
C256	GenBank Accession No. U31230, Ankri, S. et al, "Mutations in the Corynebacterium glutamicum proline biosynthetic pathway: a natural bypass of th proA step," <i>J. Bacteriol.</i> , Vol. 178(15):4412-4419 (1996), 08/02/96
C257	GenBank Accession No. U31281 for Two new members of the bio B superfamily: cloning, sequencing and expression of bio B genes of Methylobacillus flagellatum and Corynebacterium glutamicum, Serebriiskii, I.G. et al, 11/21/96
C258	GenBank Accession No. U35023 for A Corynebacterium glutamicum gene encoding a two-domain protein similar to biotin carboxylases and biotin-carboxyl-carrier proteins, Jager, W. et al, 01/16/97
C259	GenBank Accession No. U43535 for A Corynebacterium glutamicum gene conferring multidrug resistance in the heterologous host Escherichia coli, Jager, W. et al, 04/09/97
C260	GenBank Accession No. U43536 for Corynebacterium glutamicum, Jaeger, W. et al, 03/13/97
C261	GenBank Accession No. U53587 for Utilization of IS1207 for insertional mutagenesis in Corynebacteria, Bonamy, C. et al, 05/06/96
C262	GenBank Accession No. U89648 for Corynebacterium glutamicum, Kim, S.Y. et al, 03/30/99
C263	GenBank Accession No. X04960 for complete nucleotide and deduced amino acid sequences of the Brevibacterium lactofermentum tryptophan operon, Matsui, K. et al, 04/18/05
C264	GenBank Accession No. X07563 for Nucleotide sequence of the lysA gene of Corynebacterium glutamicum and possible mechanisms for modulation of its expression, Yeh, P. et al, 04/18/05
C265	GenBank Accession No. X14234 for The phosphoenolpyruvate carboxylase gene of Corynebacterium glutamicum: molecular cloning, nucleotide sequence, and expression, Eikmanns, B.J. et al, 04/18/05
C266	GenBank Accession No. X17313 for Molecular cloning, nucleotide sequence and fine- structural analysis of teh Corynebacterium glutamicum fda gene: structural comparison of C. glutamicum fructose-1, 6-biphosphate aldolase to class I and Class II aldolases, von der Osten, C.H. et al, 04/18/05
C267	GenBank Accession No. X53993 for Nucleotide sequence of the dapA gene from Corynebacterium glutamicum, Bonnassie, S. et al, 04/18/05
C268	GenBank Accession No. X54223 for DNA sequence homology between att B-related sites of corynebacterium diphtheriae, Corynebacterium ulcerans, Corynebacterium glutamicum, and the attP site of lambda-corynephage, Cianciotto, N. et al, 12/17/92
C269	GenBank Accession No. X54740 for Nucleotide sequence and organization of the upstream region of the Corynebacterium glutamicum lysA gene, Marcel, T. et al, 04/18/05
C270	GenBank Accession No. X55994 for Nucleotide sequence of the Corynebacterium glutamicum
10-10	

Substitute for form 1449A/B/PTO				Complete if Known		
				Application Number	10/627476-Conf. #2236	
INF	ORMATIC	ON DIS	SCLOSURE	Filing Date	July 25, 2003	
STATEMENT BY APPLICANT			PPLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many	sheets as	necessary)	Examiner Name	John S. Brusca	
neet	15	of	19	Attorney Docket Number	BGI-125CPCN	

	trpE gene, Heery, D.M. et al, 04/18/05
C271	GenBank Accession No. X56037 for The molecular structure of the Corynebacterium
	glutamicum threonine synthase gene, Han, K.S. et al, 04/18/05
C272	GenBank Accession No. X56075 for DNA sequence homology between att B-related sites of
	corynebacterium diphtheriae, Corynebacterium ulcerans, Corynebacterium glutamicum, adn
	the attP site of lambda-corynephage, Cianciotto, N. et al, 12/17/92
C273	GenBank Accession No. X57226 for Aspartokinase genes lysC alpha and lysC beta overlap
02.0	and are adjacent to the aspartate beta-semialdehyde dehydrogenase gene asd
	Corynebacterium glutamicum, Kalinowski, J. et al, 04/18/05
C274	GenBank Accession No. X59403 for Identification, sequence analysis, and expression of a
02/4	Corynebacterium glutamicum gene cluster encoding the three glycolytic enzymes
	glyceraldehyde-3-phosphate dehydrogenase, 3-phosphoglycerate kinase, and
0075	triosephosphate isomerase, Eikmanns, B.J. et al, 04/18/05
C275	GenBank Accession No. X59404, Bormann, E.R. et al, "Molecular analysis of the
	Corynebacterium glutamicum gbh gene encoding glutamate dehydrogenase," Mol. Microbiol.,
	Vol. 6(3):317-326 (1992), 04/18/05
C276	GenBank Accession No. X60312 for Molecular analysis of the Corynebacterium glutamicum
	lysl gene involved in lysine uptake, Seep-Feldhaus, A.H. et al, 01/30/92
C277	GenBank Accession No. X66078 for Cloning and nucleotide sequence of the csp1 gene
	encoding PS1, one of the two major secreted proteins of Corynebacterium glutamicum: the
	deduced N-terminal region of PS1 is similar to the Mycobacterium antigen 85 complex, Joliff,
	G. et al, 06/30/93
C278	GenBank Accession No. X66112, Eikmanns, B.J. et al, "Nucleotide sequence, expression and
	transcriptional analysis of the Corynebacterium glutamicum gltA gene encoding citrate
	synthase," <i>Microbiology,</i> Vol. 140(pt. 8):1817-1828 (1994), 04/18/05
C279	GenBank Accession No. X67737 for Corynebacterium glutamicum, Eikmanns, B.J. et al,
	04/18/05
C280	GenBank Accession No. X69103 for Characterization of the cspB gene encoding PS2, an
	ordered surface-layer protein in Corynebacterium glutamicum, Peyret, J.L. et al, 09/09/04
C281	GenBank Accession No. X69104 for Identification of IS1206, a Corynebacterium glutamicum
	IS3-related insertion sequence and phylogenetic analysis, Bonamy, C. et al, 09/09/04
C282	GenBank Accession No. X70584 for In vivo comparison of zidovudine resistance mutations in
	blood and CSF of HIV-1-infected patients, Wildemann, B. et al, 08/06/95
C283	GenBank Accession No. X70959 for Leucine synthesis in Corynebacterium glutamicum:
0200	enzyme activities, structure of leuA, and effect of leuA inactivation on lysine synthesis, Patek,
	M. et al, 09/09/04
C284	GenBank Accession No. X71489 for Cloning, sequence analysis, expression, and inactivation
0204	of the Corynebacterium glutamicum icd gene encoding isocitrate dehydrogenase and
	biochemical characterization of the enzyme, Eikmanns, B.J. et al, 04/18/05
C285	
C265	GenBank Accession No. X72855, Guyonvarch, A. et al, "Glutamate dehydrogenase (gdhA)
	gene." 04/18/05
C286	GenBank Accession No. X75083 for A sequence from a tryptophan-hyperproducing strain of
	Corynebacterium glutamicum encoding resistance to 5-methyltryptophan, Heery, D.M. et al,
	08/18/94
C287	GenBank Accession No. X75085 for Construction and characterization of recA mutant strains
	of Corynebacterium glutamicum and Brevibacterium lactofermentum, Fitzpatrick, R. et al,
	04/18/05
C288	GenBank Accession No. X75504 for Characterization of the isocitrate lyase gene from
	Corynebacterium glutamicum and biochemical analysis of the enzyme, Reinscheid, D.J. et al,
	04/18/05

Examiner Signature Date Considered

Substitute for form 1449A/B/PTO				Complete if Known		
				Application Number	10/627476-Conf. #2236	
00	VFORMATIO	N DISC	CLOSURE	Filing Date	July 25, 2003	
	TATEMENT	BY AF	PLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many s	heets as ne	cessary)	Examiner Name	John S. Brusca	
Sheet	16	of	19	Attorney Docket Number	BGI-125CPCN	

	comparative sequence analysis of elongation factor Tu and ATP-synthase beta-subunit genes, Ludwig, W. et al, 10/27/94
C290	GenBank Accession No. X77034 for Phylogenetic relationships of Bacteria based on
	comparative sequence analysis of elongation factor Tu and ATP-snythase beta-subunit genes, Ludwig, W. et al, 04/18/05
C291	GenBank Accession No. X77384 for Nucleotide sequence of a recA gene from Corynebacterium glutamicum, Billman-Jacobe, H. et al, 04/18/05
C292	GenBank Accession No. X78491 for Malate synthase from Corynebacterium glutamicum: sequence analysis of the gene and biochemical characterization of the enzyme, Reinscheid, D.J. et al, 04/18/05
C293	GenBank Accession No. X80629 for Phylogenetic analysis of the genera Rhodococcus and Norcardia and evidence for the evolutionary origin of the genus Nocardia from within the radiation of Rhodococcus species, Rainey, F.A. et al, 04/01/04
C294	GenBank Accession No. X81191 for Structure of the gluABCD cluster encoding the glutamate uptake system of Corynebacterium glutamicum, Kronemeyer, W. et al, 04/18/05
C295	GenBank Accession No. X81379, Wehrmann, A. et al, "Analysis of different DNA fragments of Corynebacterium glutamicum complementing dapE of Escherichia coli," <i>Microbiology</i> , Vol.140(pt. 12):3349-3356 (1994), 02/25/03
C296	GenBank Accession No. X82061 for Phylogeny of the genus Corynebacterium deduced from analysis of small-subunit ribosomal DNA sequences, Ruimy, R. et al, 11/10/95
C297	GenBank Accession No. X82928 for Multicopy suppression by asd gene and osmotic stress- dependent complementation by heterologous proA in proA mutants, Serebrijski, I. et al, 04/18/05
C298	GenBank Accession No. X82929 for Multicopy suppression by asd gene and osmotic stress- dependent complementation by heterologous proA in proA mutants, Serebrijski, I. et al, 04/18/05
C299	GenBank Accession No. X84257 for Phylogenetic analysis of the genus Corynebacterium based on 16S rRNA gene sequences, Pascual, C. et al, 01/09/04
C300	GenBank Accession No. X85965, Wehrmann, A. et al, "Functional analysis of sequences adjacent to dapE of Corynebacterium glutamicum reveals the presence of aroP, which encodes the aromatic amino acid transporter," <i>J. Bacteriol.</i> , Vol. 177(20):5991-5993 (1995), 11/30/97
C301	GenBank Accession No. X86157, Sakanyan, V. et al, "Genes and enzymes of the acetyl cycle of arginine biosynthesis in Corynebacterium glutamicum: enzyme evolution in the early steps of the arginine pathway," <i>Microbiology,</i> Vol. 142(Pt. 1):99-108 (1996), 04/18/05
C302	GenBank Accession No. X86780 for The biosynthetic gene cluster for the polyketide immunosuppressant rapamycin, Schwecke, T. et al, 04/18/05
C303	GenBank Accession No. X89084 for Cloning, sequence analysis, expression and inactivation of the Corynebacterium glutamicum pta-ack operon encoding phosphotransacetylase and acetate kinase, Reinscheid, D.J. et al, 04/18/05
C304	GenBank Accession No. X89850 for Genetic characterization of site-specific integration functions of phi AAU2 infecting 'Arthrobacter aureus' C70, Le Marrec, C. et al, 08/08/96
C305	GenBank Accession No. X90356 for Promoters from Corynebacterium glutamicum: cloning, molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
C306	GenBank Accession No. X90357 for Promoters from Corynebacterium glutamicum: cloning, molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
C307	GenBank Accession No. X90358 for Promoters from Corynebacterium glutamicum: cloning, molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
C308	GenBank Accession No. X90359 for Promoters from Corynebacterium glutamicum: doning, molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
C309	GenBank Accession No. X90360 for Promoters from Corynebacterium glutamicum: cloning,
miner	Date

Substitute for form 1449A/B/PTO				Complete if Known		
				Application Number	10/627476-Conf. #2236	
	IFORMATION	I DIS	SCLOSURE	Filing Date	July 25, 2003	
S	TATEMENT B	BY A	APPLICANT	First Named Inventor	Markus POMPEJUS	
				Art Unit	1631	
	(Use as many sh	eets as	necessary)	Examiner Name	John S. Brusca	
Sheet	17	of	19	Attorney Docket Number	BGI-125CPCN	

-		molecular analysis and search for a consensus matif. Datak M. et al. 44/04/06
├	C240	molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
1 1	C310	GenBank Accession No. X90361 for Promoters from Corynebacterium glutamicum: cloning,
<u> </u>	0044	molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
'	C311	GenBank Accession No. X90362 for Promoters from Corynebacterium glutamicum: cloning,
	0040	molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
1 19	C312	GenBank Accession No. X90363 for Promoters from Corynebacterium glutamicum: cloning,
ļ		molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
	C313	GenBank Accession No. X90364 for Promoters from Corynebacterium glutamicum: cloning,
		molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
	C314	GenBank Accession No. X90365 for Promoters from Corynebacterium glutamicum: cloning,
		molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
i 19	C315	GenBank Accession No. X90366 for Promoters from Corynebacterium glutamicum: cloning,
		molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
(C316	GenBank Accession No. X90367 for Promoters from Corynebacterium glutamicum: cloning,
		molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
	C317	GenBank Accession No. X90368 for Promoters from Corynebacterium glutamicum: cloning,
l		molecular analysis and search for a consensus motif, Patek, M. et al, 11/04/96
	C318	GenBank Accession No. X93513, Siewe, R.M. et al, "Functional and genetic characterization
1		of the (methyl)ammonium uptake carrier of Corynebacterium glutamicum," J. Biol. Chem., Vol.
		271(10):5398-5403 (1996), 05/29/96
	C319	GenBank Accession No. X93514 for Isolation, characterization, and expression of the
		Corynebacterium glutamicum betP gene, encoding the transport system for the compatible
		solute glycine betaine, Peter, H. et al, 09/08/97
	C320	GenBank Accession No. X95649 for Identification and transcriptional analysis of the dapB-
1		ORF2-dapA-ORF4 operon of Corynebacterium glutamicum, encoding two enzymes involved in
		L-lysine synthesis, Patek, M. et al, 12/21/00
	C321	GenBank Accession No. X96471 for A new type of transporter with a new type of cellular
		function: L-lysine export from Corynebacterium glutamicum, Vrljic, M. et al, 04/18/05
	C322	GenBank Accession No. X96580, Sahm, H. et al, "D-Pantothenate synthesis in
	J J	Corynebacterium glutamicum and use of panBC and genes encoding L-valine synthesis for D-
		pantothenate overproduction," <i>Appl. Environ. Microbiol.</i> , Vol. 65(5):1973-1979 (1999),
		04/18/05
- (C323	GenBank Accession No. X96962 for Utilisation of IS1207 for insertional mutagenesis in
	0	Corynbacteria, Bonamy, C. et al, 07/07/02
1	C324	GenBank Accession No. X99289 for Cloning, sequencing and expression of the gene
	OOL ,	encoding elongation factor P in the amino-acid producer Brevibacterium lactofermentum
		(Corynebacterium glutamicum ATCC 13869), Ramos, A. et al, 09/09/04
l - 1	C325	GenBank Accession No. Y00140 for Nucleotide sequence of the homoserine kinase (thr B)
I '	OOLO	gene of Brevibacterium lactofermentum, Mateos, L.M. et al, 09/12/93
l	C326	GenBank Accession No. Y00151 for Nucleotide sequence of the meso-diaminopimelate D-
	سک	dehydrogenase gene from Corynebacterium glutamicum, Ishino, S. et al, 09/12/93
 	C327	
 	WZ1	GenBank Accession No. Y00476 for Nucleotide sequence of the homoserine dehydrogenase
 	C328	(thr A) gene of Brevibacterium lactofermentum, Mateos, L.M. et al, 05/05/93
	W20	GenBank Accession No. Y00546 for Nucleotide sequence and fine structural analysis of the
}	0220	Corynebacterium glutamicum hom-thrB operon, Peoples, O.P. et al, 09/12/93
	C329	GenBank Accession No. Y08964 for Identification, characterization, and chromosomal
		organization of the ftsZ gene from Brevibacterium lactofermentum, Honrubia, M.P. et al,
	0000	04/18/05
	C330	GenBank Accession No. Y09163 for Isolation of the putP gene of Corynebacterium
		Iglutamicum and characterization of a low-affinity uptake system for compatible solutes, Peter,
		H. et al, 09/08/97
Examiner		Date
Signature	<u></u>	Considered

BGI-125CPCN

Considered

PTO/SB/08a/b (07-05)

Approved for use through 07/31/2006. OMB 0651-0031

U.S. Patent and Trademark Office; U.S. DEPARTMENT OF COMMERCE

Under the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it contains a valid OMB control number. Complete if Known Substitute for form 1449A/B/PTO 10/627476-Conf. #2236 Application Number INFORMATION DISCLOSURE July 25, 2003 Filing Date STATEMENT BY APPLICANT Markus POMPEJUS First Named Inventor 1631 **Art Unit** (Use as many sheets as necessary) John S. Brusca **Examiner Name**

Attorney Docket Number

19

Sheet

Signature

18

of

C331	juniority and the state of the				
	characterization, expression and inactivation of the pyc gene, Peters-Wendisch, P.G. et al, 04/18/05				
C332	GenBank Accession No. Y09578 for Analysis of the leuB gene from Corynebacterium				
	glutamicum, Patek, M. et al, 04/18/05				
C333	GenBank Accession No. Y12472 for Site-specific integration of corynephage phi16: the				
	construction of an integration vector, Moreau, S. et al, 03/05/99				
C334	GenBank Accession No. Y12537 for Corynebacterium glutamicum is equipped with four				
	secondary carriers for compatible solutes: identification, sequencing, and characterization of the proline/glycine betaine carrier, EctP, Peter, H. et al, 11/17/98				
C335	GenBank Accession No. Y13221 for Isolation of the Corynebacterium glutamicum glnA gene encoding glutamine synthetase I, Jakoby, M. et al, 08/28/97				
C336					
	mycobacterial two-component system operon, Supply, P. et al, 04/18/05				
C337					
C338					
0556	GenBank Accession No. Y18059 for Analysis of the integration functions of phi304L: an integrase module among corynephages, Moreau, S. et al, 09/29/99				
C339					
	the upstream region of the lysA gene in Brevibacterium lactofermentum: regulation of argS-				
	lysA cluster expression by arginine, Oguiza, J.A. et al, 04/18/05				
C340					
	Brevibacterium lactofermentum encodes dihydrodipicolinate synthase, dihydrodipicolinate				
	reductase, and a third polypeptide of unknown function, Pisabarro, A. et al, 08/16/93				
C341					
	Brevibacterium lactofermentum and characterization of the encoded threonine synthase,				
	Malumbres, M. et al, 04/18/05				
C342					
	al, 11/21/94				
C343	GenBank Accession No. Z49822 for Multiple sigma factor genes in Brevibacterium				
	lactofermentum: characterization of sigA and sigB, Oguiza, J.A. et al, 04/18/05				
C344					
	of Brevibacterium lactofermentum is coupled transcriptionally to the dmdR gene, Oguiza, J.A. et al, 04/18/05				
C345	GenBank Accession No. Z49824 for Multiple sigma factor genes in Brevibacterium				
	lactofermentum: characterization of sigA and sigB, Oguiza, J.A. et al, 04/18/05				
C346					
	present in the genome of Brevibacterium lactofermentum ATCC 13869, Correia, A. et al, 07/07/02				
C347	GenBank Accession No. Z77162 for Deciphering the biology of Mycobacterium tuberculosis				
	from the complete genome sequence, Cole, S.T. et al, 09/02/02				
C348					
	from the complete genome sequence, Cole, S.T. et al, 09/02/02				
C349					
	tuberculosis from the complete genome sequence," <i>Nature</i> , Vol. 393(6685):537-544 (1998),				
	09/02/02				
C350	GenBank Accession No. Z98209, Cole, S.T. et al, "Deciphering the biology of Mycobacterium tuberculosis from the complete genome sequence," <i>Nature</i> , Vol. 393(6685):537-544 (1998),				
	08/03/01				
C351	GenBank Accession No. Z83866 for Deciphering the biology of Mycobacterium tuberculosis from the complete genome sequence, Cole, S.T. et al, 09/02/02				

PTO/SB/08a/b (07-05)

Approved for use through 07/31/2006. OMB 0651-0031

U.S. Patent and Trademark Office; U.S. DEPARTMENT OF COMMERCE

Under the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it contains a valid OMB control number. Complete if Known Substitute for form 1449A/B/PTO 10/627476-Conf. #2236 **Application Number INFORMATION DISCLOSURE** July 25, 2003 Filing Date STATEMENT BY APPLICANT Markus POMPEJUS First Named Inventor

_		- , ,				
				Art Unit	1631	•
	(Use as many she	eets as	s necessary)	Examiner Name	John S. Brusca	
Sheet	19	of	19	Attorney Docket Number	BGI-125CPCN	•

C352	GenBank Accession No. Z95388, Cole, S.T. et al, "Deciphering the biology of Mycobacterium tuberculosis from the complete genome sequence," <i>Nature</i> , Vol. 396(6685):537-544 (1998) Feb. 10, 1999	
C353	* Eggleling, L. et al, "Transport Mutants and Transport Genes of Corynebacterium glutamicum," Annals of the New York Academy of Sciences, Vol. 782:191-201 (1996)	
C354	Siewe, Ruth M. et al, "Functional and Genetic Characterization of the (Methyl)ammonium Uptake Carrier of Corynebacterium glutamicum," The Journal of Biological Chemistry, Vol. 271(10):5398-5403 (1996)	
C355	* GenBank Accession No. A89996, Kuroda et al, "Probable Ammonium Transporter nrgA, Protein Information Resource Database," May 2001	
C356	Lee, J.K. et al. "Nucleotide sequence of the gene encoding the Corynebacterium glutamicum mannose enzyme II and analyses of the deduced protein sequence," FEMS Microbiol. Lett., 119(1-2):137-145 (1994)	
C357	Schäfer, A. et al. "The Corynebacterium glutamicum cgllM gene encoding a 5-cytosine in an McrBC-deficient Escherichia coli strain," <i>Gene</i> , 203(2):95-101 (1997)	

^{*}EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.

Examiner	Data	
	Jaic	4
Signature	Considered	,

^{&#}x27;Applicant's unique citation designation number (optional). ²Applicant is to place a check mark here if English language Translation is attached.

cation No. (if known): 10/627476

Attorney Docket No.: BGI-125CPCN

Certificate of Mailing under 37 CFR 1.8

I hereby certify that this correspondence is being deposited with the United States Postal Service with sufficient postage as first class mail in an envelope addressed to:

> MS Amendment **Commissioner for Patents** P.O. Box 1450 Alexandria, VA 22313-1450

on	September 28, 2006
	Date

Signature Maria Laccotripe Zacharakis, Ph.D., J.D.

Typed or printed name of person signing Certificate

56,266

(617) 227-7400

Registration Number, if applicable

Telephone Number

Each paper must have its own certificate of mailing, or this certificate must identify Note: each submitted paper.

Transmittal (1 page) Fee Transmittal (1 page, in duplicate) Information Disclosure Statement (2 pages) PTO form SB/08 (388 References) (19 pages) This Certificate of Mailing (1 page)

Copies of Three Hundred and Eighty Two (382) References

Return Receipt Postcard